

GenCore version 6.3  
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OM nucleic - nucleic search, using sw model

Run on: June 26, 2009, 03:09:48 ; Search time 1395 Seconds  
(without alignments)  
13572.847 Million cell

updates/sec

Title: US-09-300-482-619  
Perfect score: 300  
Sequence: 1  
gatttattgacaacaccgat.....ttcagaaatgtctgctgtgg 300

Scoring table: IDENTITY\_NUC  
Gapop 10.0 , Gapext 1.0

Searched: 111716937 seqs, 31560637849 residues

Total number of hits satisfying chosen parameters: 223433874

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : Pending\_Patents\_NA\_Main:\*  
1: /ABSS/Data/CRF/ptodata/1/pna/PCTUSA\_COMB.seq:\*  
2: /ABSS/Data/CRF/ptodata/1/pna/PCTUSB\_COMB.seq:\*  
3: /ABSS/Data/CRF/ptodata/1/pna/PCTUSC\_COMB.seq:\*  
4: /ABSS/Data/CRF/ptodata/1/pna/PCTUSD\_COMB.seq:\*  
5: /ABSS/Data/CRF/ptodata/1/pna/US075\_COMB.seq:\*  
6: /ABSS/Data/CRF/ptodata/1/pna/US076\_COMB.seq:\*  
7: /ABSS/Data/CRF/ptodata/1/pna/US077\_COMB.seq:\*  
8: /ABSS/Data/CRF/ptodata/1/pna/US078\_COMB.seq:\*  
9: /ABSS/Data/CRF/ptodata/1/pna/US079\_COMB.seq:\*  
10: /ABSS/Data/CRF/ptodata/1/pna/US080\_COMB.seq:\*  
11: /ABSS/Data/CRF/ptodata/1/pna/US081\_COMB.seq:\*  
12: /ABSS/Data/CRF/ptodata/1/pna/US082\_COMB.seq:\*  
13: /ABSS/Data/CRF/ptodata/1/pna/US083\_COMB.seq:\*  
14: /ABSS/Data/CRF/ptodata/1/pna/US084\_COMB.seq:\*  
15: /ABSS/Data/CRF/ptodata/1/pna/US085\_COMB.seq:\*